

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5HYC2	K2026_HUMAN	Homo sapiens	Uncharacterized protein KIAA2026	11.159387	S919;T1457;T1460;S1480;T1554;T1761;S1765;T1944;T2000	NaN	35254053;30379171;38253038;31492838;34019948;30059200;35289036	MSVPGTPGAMEPAGEEERPPPAEAG EDDEEEVAAAAQTSGPAHGRSASSL EDADDQEEEMEAMVIGGGCCCKEQE LTYELQOQYRILGEFLQEKHRGLTAP FLQPLGGVATAEEVEAEGPRSGGRG GRAFPQPGQGMCLLQMEEFASG QYGGITEFVADFRLMLETCYRLHGV DHWISKQGGKLEMMLEQKLALLSR HLREKTTIAVTSRGGYGLEDEKGTAC TSTRRRSTPRSLAGLTSGVFESIMVQ VLRQEEQLRAKEEKRLREQERKEAE EASQKEIEEWERKLLAQAAPTCMET MWEIPAIGHFLCLAQQILNLPEIVFY ELERCLLMPOCNAFLSKIMTSLLSLSP PHRRPTLHRRPTLPYRTWEAALRQK VQQWYTAGVQATENPDNCAEKLGLC PQFFKVLGEVNPLEEKPFHELPHYQ KVVLLKGLCDFVYETQKEVQDAVLG QPIHECREVILGYDYLENAYVHFPQF CGADVRIYKQRPFOAPEFPPIPIKIQR VPRIKLEKLCQDQVSTSNGEHRCSR DSLPSFKEQENNFDPACCPAKMI LDNHDISVEMGVKSNYEIRIRRPCEI KKTDCCKENLEKPRSPGEVTGFGEPE LSPGEIRFIENQEKYGEASRIKIEPSP LKENTLKSQIHVNGSHSDHPEINC HKVVRDILLEQSLQSHKLLKLTMR AKKKKKKKKLDVLENNLQKRE GLHSLAFKSYKPEIQNKLLIHKKAK HKKHKS GKSVSKAITKKRKTVIKS PTVPEFQLICTNLDELRELITKIENEL KDLNLSRKKSGKWYHRRQAVKELH STLIRLLNELLPWEPKLMKAFQRNR SRLKKDYDDFRROPDHDFTNRELW TTDEGEGDLGKDSPKGEISKSIDSTE PLDILEKDHFDSDDMKLEIDFPMA RSKLLKELPSKDLPKTLLKTLKRQS KQTDYVDDSTKELSPRKKAKLSTNE TTVENLESVDVQIDCFSESKHTEPSFP ESFASLDSVPVSTLQKGTKPIQALLA KNIGNKVTLTNQLPPSTGRNALAVE KPVLSPEASPIKALTCHTNTKGPL QMVKMPCGQWLPIDLQNSSVKIQV QPMVDPKTGEKIMQQVLILPKNFVI QHKEGKAVAKEVPPLQKQKTEQHC SSFQTTNINSSLASVFNNSPGTVST QLPNTAFNKTTTTPLSNISSARQPLS PVTSVSNLLTPSVKTSQSEAGKAKN AVSAATFSLPSASPTISSSTGQPLSSTT TLNGSTNPGSSFNCFAQQTADSSEA KQELKTVCIQDSQSILVRTRGGNTGV VKVQTNPDQNSPNTVSSSSVFTFAP QLQAFVLPKSTTSSSAFSPVAGTTTT SSLSPFSQTPSVSIPASFAPSMGKN LKLTLGHTTGSGDLGHVIDKTSHP SSPLKSSICSTLLPSTSSSVVISIS AANFGQNNANIHTPTKQQQVDYIT KSPVTRSEATAATNGDVISGTPVQK LMLVSAPSILSSNGTAINMTPALTS

								<p>TGVSAQKLVFINAPVPSGTSTPTLVA          ESLKQTLPPPLHKAYVKTPEQPQIVLI          PSTVGTPIKINSSPAVSQIKDVKIGLN          IGQAIVNTSGTVPAIPSINILQNVTPK          GEDKSSKGYILPLSTSGNSVPVSSNF          VSQNITPVNESVVSSARAVNVLSVTG          ANLSLGSFPVTSASASAGAQPVLVS          GNDTSSRIMPILSNRLCSSLGNTVA          ISTVKTGHLASSVLISTTQPVVSPKCL          TSALQIPVTVALPTTATSPKIINTVP          HSAAVPGATRSVSISKRSRTSLQFH          SPGISTTVPTNVNTNKPQELSSLST          SPGKITNTSNFASLPNQALVKTPSY          SSAPGGTTIHTASAPSNVTSLVGSQF          SEPCIQQKIVINTSTPLAPGTQIMING          TRFIVPPQGLGAGSHVLLISTNPKYG          APLVLNSGQGIQSTPIDNSAQKITLA          SNNLSGQPLQHLRSPTKFINSEFG          NASSIPTVHTSPQLINTTAKVPVPPPV          PTVSLTsvIKSPATLLAKTSLVSAICP          SNPPLPSSTSVFHLDPVVKLLVSPE          GAILNTINTPASKVSSLSPSLSQIVVS          ASRSPASVFPQSSGLEKPDRAAS</p>
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